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<b>TRANSMITTAL FORM</b> (to be used for all correspondence after initial filing)	<b>Application Number</b>	09/441,318	
	<b>Filing Date</b>	November 16, 1999	
	<b>First Named Inventor</b>	Conklin et al	
	<b>Group Art Unit</b>	1638	
	<b>Examiner Name</b>	Kubelik, A.	
<b>Total Number of Pages in This Submission</b>	8	<b>Attorney Docket Number</b>	BTI-41

**ENCLOSURES (check all that apply)**

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**SIGNATURE OF APPLICANT, ATTORNEY, OR AGENT**

<b>Firm or Individual name</b>	Brown & Michaels, PC
<b>Signature</b>	Thomas T. Agnello
<b>Date</b>	6/13/01

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June 13, 2001

In re Application of: Conklin *et al.*  
Serial No. 09/441,318  
Filed: November 16, 1999  
For: TRANSGENIC PLANT WITH INCREASED EXPRESSION OF GDP-MANNOSE PYROPHOSPHORYLASE  
Examiner: Kubelik, Anne R.  
Art Unit: 1638  
Attorney Docket No.: BTI-41

**PRELIMINARY AMENDMENT**

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Please amend the above-identified application as follows:

**In the Specification:**

Please delete the third paragraph on page 5, at lines 6-8, which refers to Fig. 5.

Please replace the second complete paragraph at page 7, lines 16-22, with the following replacement paragraph:

To test the hypothesis that *vtc1-1* and *vtc1-2* harbor mutations in the GDP-mannose pyrophosphorylase gene, the potential for mutations in the pyrophosphorylase genomic sequence derived from each of these mutant alleles was examined. The sequences of both *vtc1-1* and *vtc1-2* contain the identical single cytosine to thymine point mutation at position +64 relative to the first base of the presumed initiator methionine (Figure 3D). This predicted missense mutation would convert a highly conserved proline to a serine at amino acid 22 in the GDP-mannose pyrophosphorylase amino acid sequence.

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*Judy H. Barron*  
Judy H. Barron